Regex

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Github link:

https://github.com/amyrewells/biol432_As01

Group Members

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Load packages

```r
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##     filter, lag

## The following objects are masked from 'package:base':
##
##     intersect, setdiff, setequal, union
```

1

```r
MData<- read.csv("./measurements.csv")
```

2

```r
MData<- MData %>%
  mutate(Sp = sub("(\\w)\\w+", "\\1.", Species_Name))
```

3

```r
MData_new<- MData %>%
  select(Sp, everything(), -Species_Name)
```

4

```r
write.csv(MData_new, "MeansShort.csv", row.names = FALSE)
```